

5914 089

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FIG. 1A

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241?

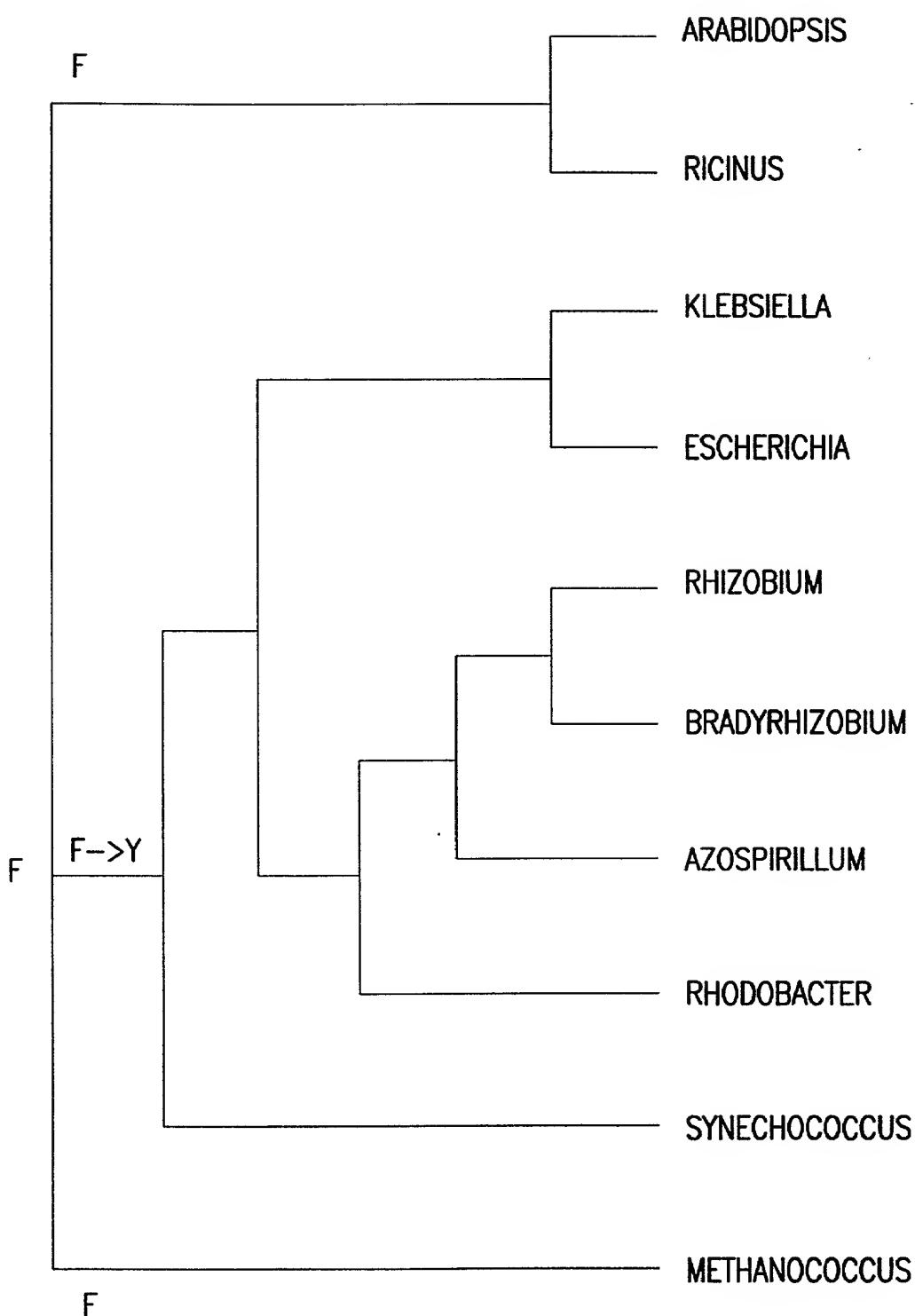


FIG. 1B

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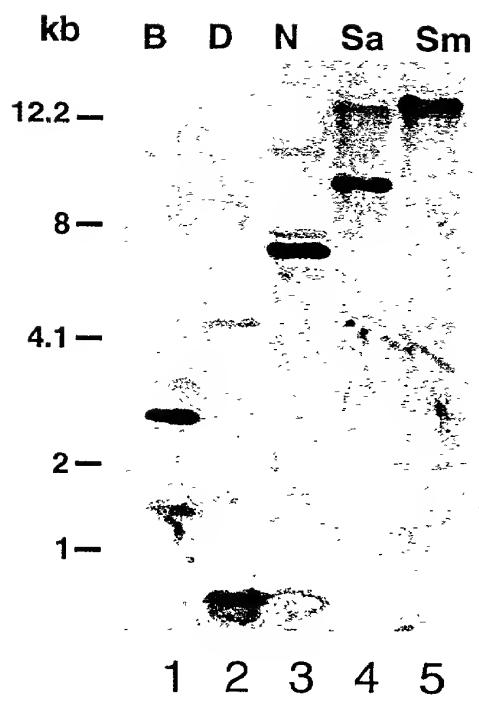


FIG.2

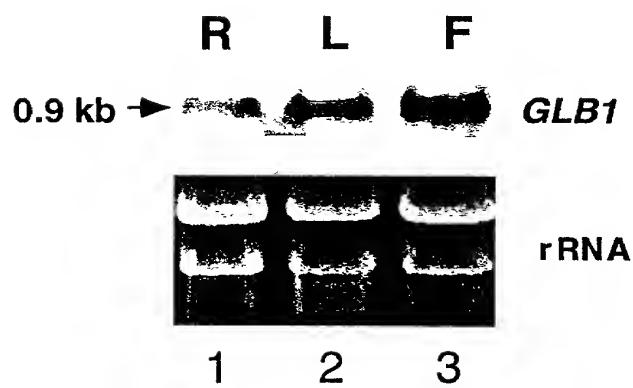


FIG.3

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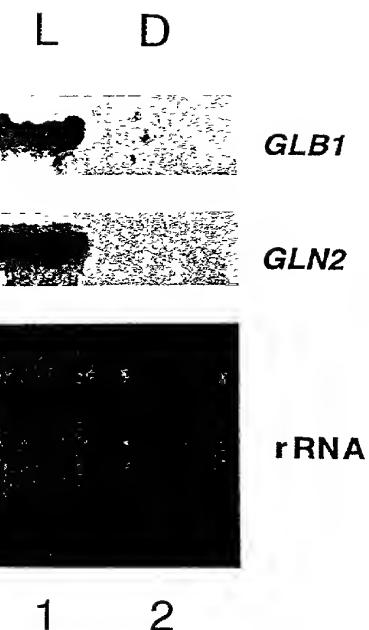


FIG.4A

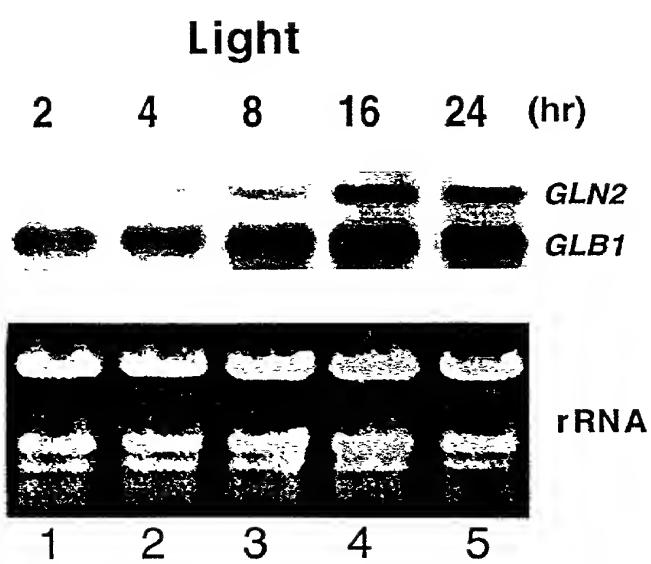


FIG.4B

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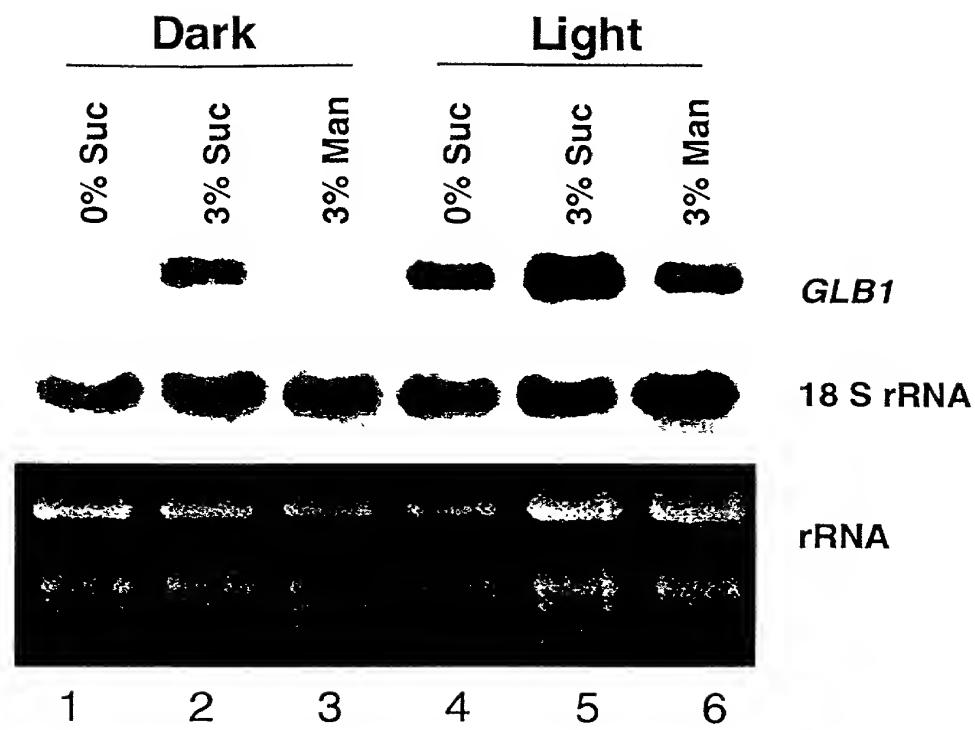


FIG.5

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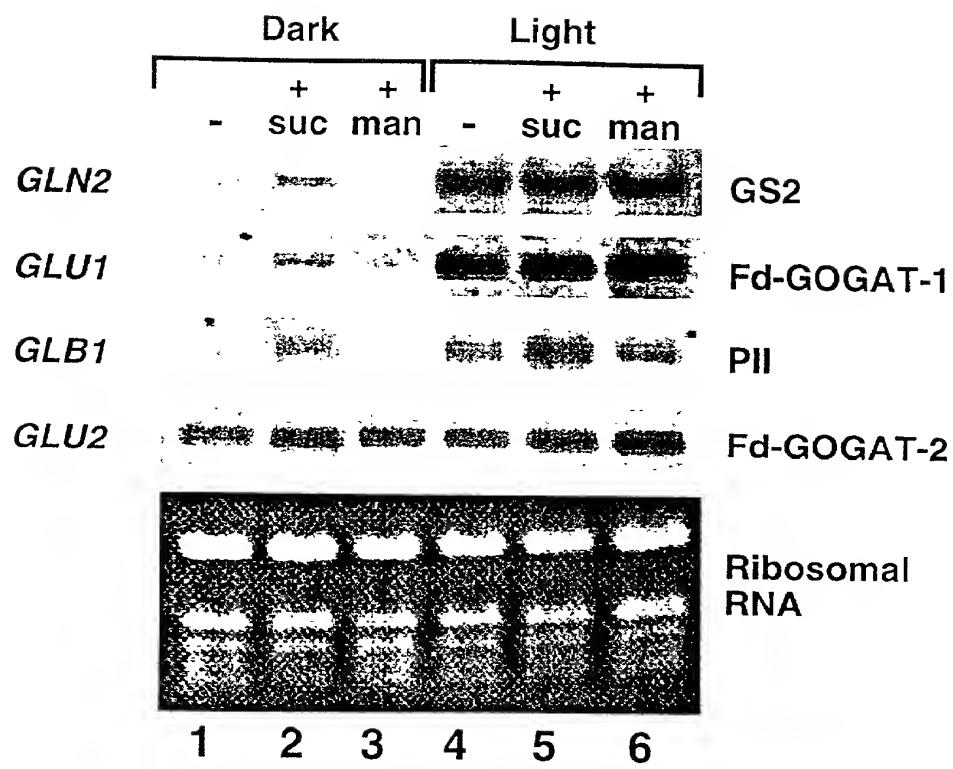


FIG. 6

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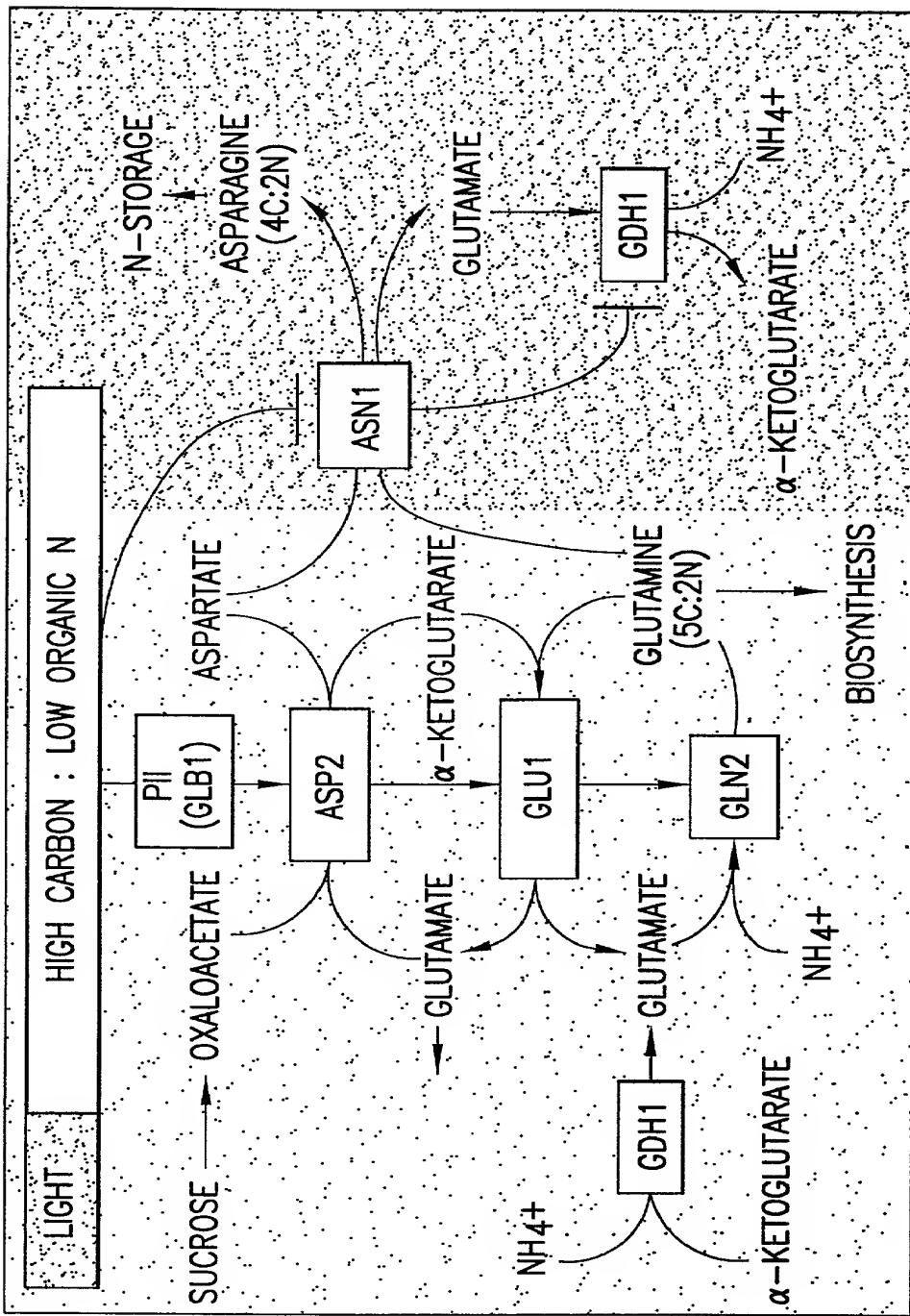


FIG. 7

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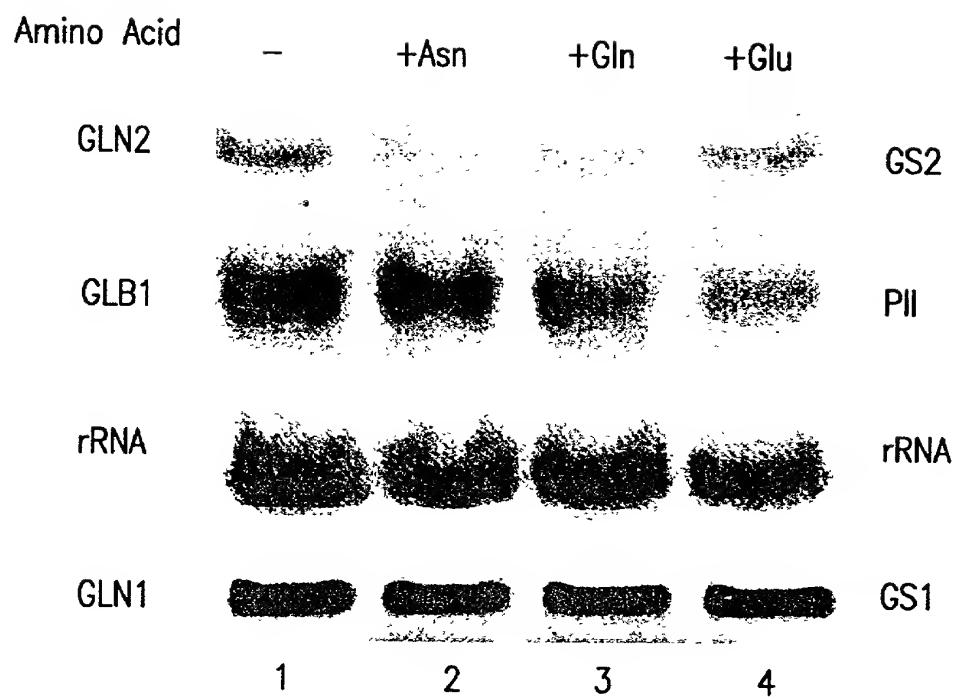


FIG. 8

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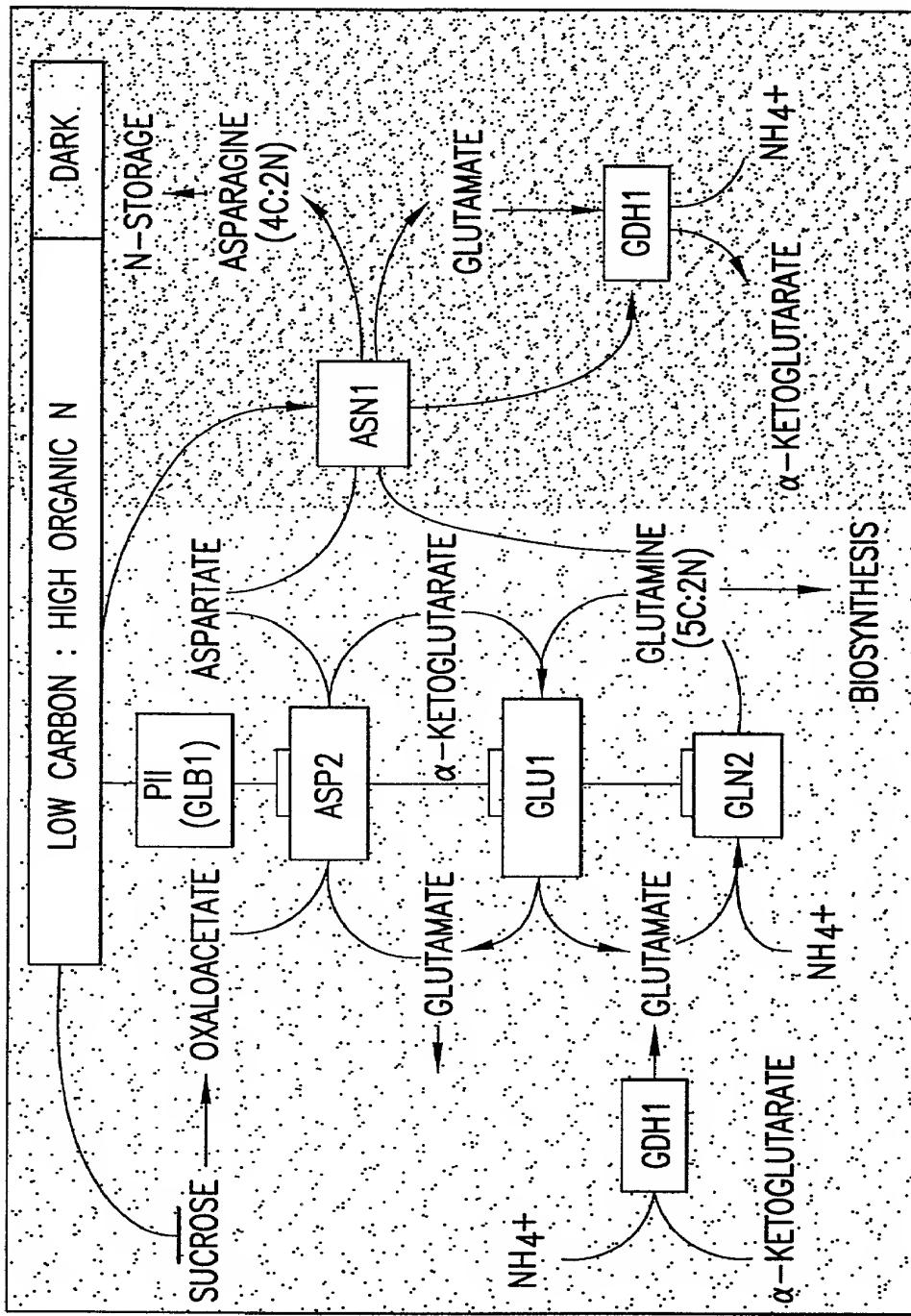


FIG. 9

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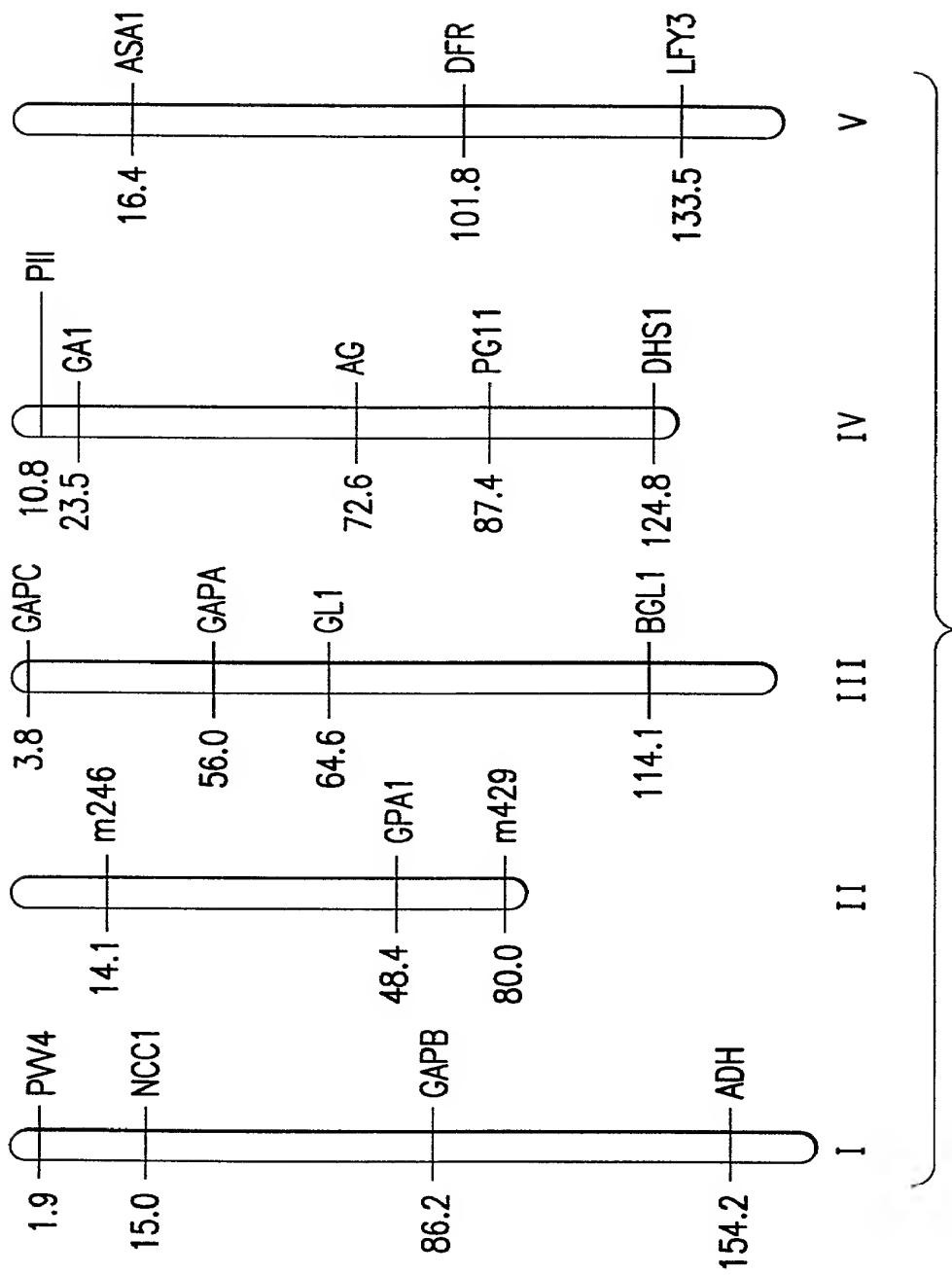


FIG. 10

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1 ctgaaagtgttaaaaaaaaaaaactagaa tcatggcggtcaatgacg
51 aaacccatct caataacttc tctcggtttc tattctgatc gaaagaacat
101 tgctttctct gattgcattt cgatttggtc tggattcaga cattcccgac
151 catcttgctt cgatttggtc acaaagtcac cgagtaataa cagtcgttt
201 ttacctgtcg tttagtgccta aatatcttct gattatattt cagactcgaa
251 attttacaag gtggaagcaa ttgtcagacc atggagaatc cagcaagttt
301 catcggcttt actgaaaaatc gggattcgag gtgttactgt ttctgtatgt
351 agagggtttt gtcacaagg aggttctacc gagagacacg gtggctctga
401 gttctcgaa gacaaattt gttgctaaagt taagatggaa atcgttgtta
451 agaaagacca agtggaatct gtaatcaaca caataattga aggagcaagg
501 acaggagaga ttggatgttcaagattttt gtttgcctg tgtcagatgt
551 cataagagttt aggacaggtt agcgtgggaa gaaagcagag aagatgactg
601 gtgatatgtt ttcaccgtct taggaacaaa cagagctcaa gaatggttt
651 ttttttttc atttcggtct ctagattctg cgaataataa tgaatggagt
701 ctgtgtttgg tttcatgttg aatcgatcaa gatgtgtttt taactgtaca
751 tgaattatgc agaaaacatct gtcctggttc tcagacatcg aaactctgtt
801 cctaataaaaa aaaaaaaaaaaaaaaa

FIG.12

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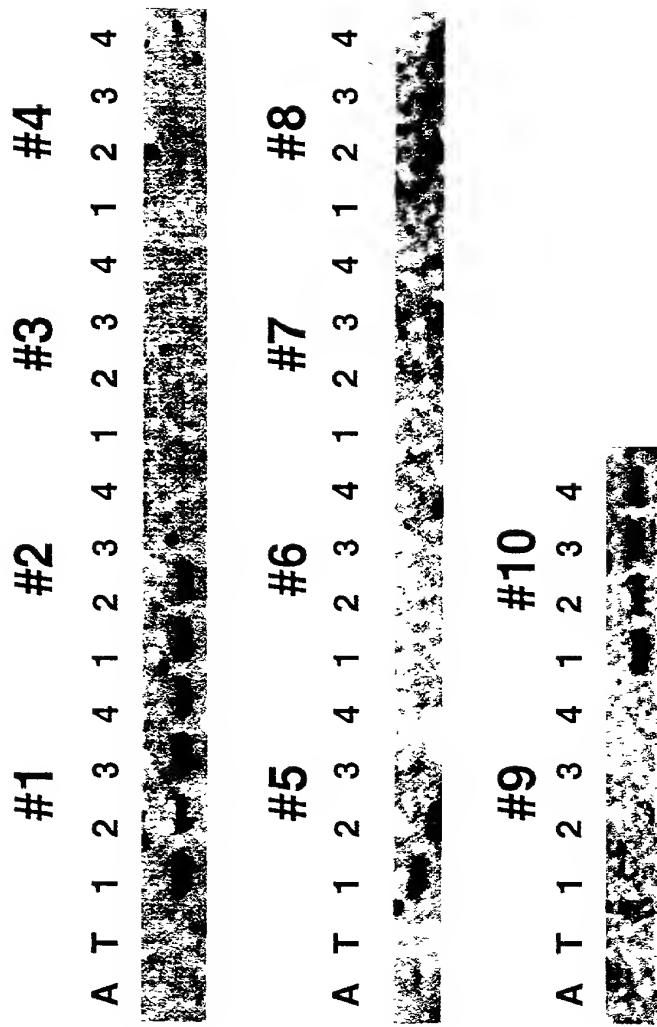


FIG. 11

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1 GCGGTGTCGG CCGCTCTAGA ACTAGTGGAT CCCCCGGGCT GCAGGAATT
51 GGCACGAGGC TACTGCGAAA CTGGGCTTGC TCACTCCTCT TCATTCTAAT
101 AACATCAAGA AAGAATTCCC TGTTTTGAT TTCAGTTGT TTTGTCCAGA
151 GCTTAGACAT TCTCGGTTTT CTCACTTTAA CACCGCGGTC AAGCGCGTAA
201 GATATGCCCC CGTCGTTCCCT GTGATTAATG CCCAAAGCTC GCCTGACTAC
251 ATTCCCTGATG CTAAATTCTA CAAAGTGGAA GCAATTCTCA GGCCCTGGCG
301 AGTCTCGCAA GTTTCCTCGG CTTTGCTAAA AATTGGTATT CGAGGGTGT
351 CTGTTCTGA TGTCGAGGT TTTGGTGCTC AAGGTGGTTC AACTGAGAGG
401 CAGGGCGGCT CAGAATTTC TGAAGACAAG TTTGTTGCTA AAGTTAAGAT
451 GGAGATCGTG GTTAGCAAAG ACCAGGTTGA GGATGTTATA GAAAAAAATCA
501 TTGAGGAGGC AAGAACTGGA GAGATTGGAG ACGGCAAGAT TTTCTTGCTG
551 CCTGTTTCAG ATGTAATAAG AGTCGCACT GGTGAGCGGG GTGATAAGGC
601 TGAGAGGATG ACAGGAGGGC GATCTGACAT GAGTACTTCT GCTTGACTGC
651 TGTGACCAGC AATATAGCAT TCAGGACTAA CTGTCCTTTG AGAAAGCCCC
701 GCCCTTATTAA GCCATTATCC AGTATAGCTT GATAATTGATTTTTGTT
751 TTCTTAACCA AAGAAACAAA GATCTTTCA TTATCCTGTT GATGATAATT
801 GAAAACGGAA GGATCGCGAA TTTGTTCAAG TGCTTGCAAG ATAAATAACA
851 AGAAGAGGAG TAATGTTAAC AAAAAAAAAA AAAAAAAAAA ACTCGAG

FIG.13